SEQUENCE LISTING

<110> Mirochnitchenki, O.
Wei, J.
Inouye,M.

<120> Ischemia Activated Protein

<130> 13257-00026

<140> tba
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<160> 11

<170> FastSEQ for Windows Version 3.0

<210> 1
<211> 744
<212> DNA
<213> Homo sapiens

<400> 1

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gcggcccccg  gcgcccggct  gttgcggctc  ccggggagcg  gggccgtgca  ggccgcgagc   180
ccggagcgcg  ccggctggac  cgaggcgctg  cgggccgccg  tggccgagct  gcgcgccggc   240
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<210> 2
<211> 100
<212> PRT
<213> Homo sapiens

<400> 2

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Ser Val Gly Leu Ser Glu Gly Pro Ala Gly Ser Arg Ser Gly Arg Leu
            20                  25                  30
Phe Arg Pro Pro Ser Pro Ala Pro Ala Ala Pro Gly Ala Arg Leu Leu
        35                  40                  45
Arg Leu Pro Gly Ser Gly Ala Val Gln Ala Ala Ser Pro Glu Arg Ala
    50                  55                  60
Gly Try Thr Glu Ala Leu Arg Ala Ala Val Ala Glu Leu Arg Ala Gly
65                  70                  75                  80
```

```
Ala Val Val Ala Val Pro Thr Asp Thr Leu Tyr Gly Leu Ala Cys Ala
                85                  90                  95
Ala Ser Cys Ser Ala Ala Leu Arg Ala Val Tyr Arg Leu Lys Gly Arg
        100                 105                 110
Ser Glu Ala Lys Pro Leu Ala Val Cys Leu Gly Arg Val Ala Asp Val
    115                 120                 125
Tyr Arg Tyr Cys Arg Val Arg Val Pro Glu Gly Leu Leu Lys Asp Leu
130                 135                 140                 145
Leu Pro Gly Pro Val Thr Leu Val Met Glu Arg Ser Glu Glu Leu Asn
                150                 155                 160
Lys Asp Leu Asn Pro Phe Thr Pro Leu Val Gly Iso Arg Iso Pro Asp
            165                 170                 175
His Ala Phe Met Gln Asp Leu Ala Gln Met Phe Glu Gly Pro Leu Ala
        180                 185                 190
Leu Thr Ser Ala Asn Leu Ser Ser Gln Ala Ser Ser Leu Asn Val Glu
    195                 200                 205
Glu Phe Gln Asp Leu Try Pro Gln Leu Ser Leu Val Iso Asp Gly Gly
210                 215                 220                 225
Gln Iso Gly Asp Gly Gln Ser Pro Glu Cys Arg Leu Gly Ser Thr Val
                230                 235                 240
Val Asp Leu Ser Val Pro Gly Lys Phe Gly Iso Iso Arg Pro Gly Cys
            245                 250                 255
Ala Leu Glu Ser Thr Thr Ala Iso Leu Gln Gln Lys Tyr Gly Leu Leu
        260                 265                 270
Pro Ser His Ala Ser Tyr Leu
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<210> 3
<211> 1387
<212> DNA
<213> Homo sapiens

<400> 3

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ggtccgcttg  ctctcactag  tgccaacctc  agctcccagg  ccagttctct  gaatgtcgag     180
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gatttagaat  ctagtgagag  cntcctctct  gggggtggtg  gcatttaagg  ttcaacccan     780
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cagcctgggc  aacatagcaa  tactccgtct  cataaaaatt  aataaataaa  aagtctcagg     960
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tccgcagcct  tttgtctatt  tatatgtgta  ccaagtgtta  aacataatta  ttattgggca    1320
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<210> 4
<211> 1048
<212> DNA
<213> Mus musculus

<400> 4

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ccggcgctgc  cgggggcccg  gctgctgcgg  cttccggaga  gcgagcccgt  ggaagccgcg    180
agccccgagc  gcgccggctg  gaccgaggcg  ctgcgggccg  ccgtggccga  gctgcgcgcc    240
ggcgccgtgg  tggcggtccc  gaccgacacg  ctctacggcc  tggcctgctc  ggcgagcagc    300
tcggcggccc  tgagttgcgt  gtaccgcctc  aaaggccgca  gcgaggccaa  gccgctggcc    360
gtgtgcctgg  gccgcgtggc  cgacgtctac  aggtactgtc  aggtgagagt  acctagggag    420
ctcctggaag  acctgttccc  aggccctgtg  accctggtga  tggagcgctc  cgaggagctc    480
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tcccaggcca  gttctctgag  tgttgaggag  ttccaagacc  tctggcctca  tttgtccctt    660
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<210> 5
<211> 149
<212> PRT
<213> Mus musculus

<400> 5

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            20                  25                  30
Leu Leu Leu Pro Pro Glu Pro Ala Pro Ala Leu Pro Gly Ala Arg Leu
        35                  40                  45
Leu Arg Leu Pro Glu Ser Glu Pro Val Glu Ala Ala Ser Pro Glu Arg
    50                  55                  60
Ala Gly Try Thr Glu Ala Leu Arg Ala Ala Val Ala Glu Leu Arg Ala
65                  70                  75                  80
Gly Ala Val Val Ala Val Pro Thr Asp Thr Leu Tyr Gly Leu Ala Cys
                85                  90                  95
Ser Ala Ser Ser Ser Ala Ala Leu Ser Cys Val Tyr Arg Leu Lys Gly
            100                 105                 110
Arg Ser Glu Ala Lys Pro Leu Ala Val Cys Leu Gly Arg Val Ala Asp
        115                 120                 125
Val Tyr Arg Tyr Cys Gln Val Arg Val Pro Arg Glu Leu Leu Glu Asp
    130                 135                 140
Leu Phe Pro Gly Pro Val Thr Leu Val Met Glu Arg Ser Glu Glu Leu
145                 150                 155                 160
Asn Lys Asp Leu Asn Pro Phe Thr Arg Leu Val Gly Iso Arg Iso Pro
                165                 170                 175
Asp His Ala Phe Met Leu Asp Leu Ala Gln Met Phe Gly Gly Pro Leu
            180                 185                 190
Ala Leu Thr Ser Ala Asn Leu Ser Ser Gln Ala Ser Ser Leu Ser Val
```

```
        195                 200                 205
Glu Glu Phe Gln Asp Leu Try Pro His Leu Ser Leu Val Iso Asp Gly
    210                 215                 220
Gly Pro Iso Gly Asp Ser Gln Ser Pro Glu Cys Arg Leu Gly Ser Thr
225                 230                 235                 240
Val Val Asp Leu Ser Val Pro Gly Lys Phe Gly Iso Iso Arg Pro Gly
                245                 250                 255
Cys Ala Leu Glu Asn Thr Thr Ser Iso Leu Gln Gln Lys Tyr Gly Leu
            260                 265                 270
Leu Pro Ser Gln Gly Ser Cys Ser
        275                 280
```

<210> 6
<211> 702
<212> DNA
<213> Bos Taurus

<400> 6

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catcctccag  gagtatgggc  tgctcccctc  acatggatcc  tgctggtgac actctggagg      600
agggaaggcc  caagggctgg  tgctggacac  tatgtgtccg  actgctggtg gttggcaagg      660
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<210> 7
<211> 126
<212> PRT
<213> B. Taurus

<400> 7

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Leu Val Gly Ile Arg Ile Pro Asp His Ala Phe Met Gln Asp Leu Val
            20                  25                  30
Gln Met Phe Gly Gly Pro Leu Ala Leu Thr Ser Ala Asn.Leu Ser Ser
        35                  40                  45
Gln Ser Ser Ser Leu Asn Val Glu Glu Phe Gln Asp Leu Trp Pro His
    50                  55                  60
Leu Ser Leu Ile Ile Gly Gly Gly Pro Ile Gly Asp Gly Gln Ser Pro
65                  70                  75                  80
Glu Cys Arg Leu Gly Ser Thr Val Val Asp Leu Ser Val Pro Gly Lys
                85                  90                  95
Phe Gly Ile Ile Arg Pro Gly Cys Ala Leu Glu Ser Thr Ser Ala Ile
            100                 105                 110
Leu Gln Glu Tyr Gly Leu Leu Pro Ser His Gly Ser Cys Trp
        115                 120                 125
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<210> 8

<211> 841
<212> DNA
<213> Rattus novartis

<400> 8

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catgctggat actatgtgtc tactacaggt tggcaaagcc tcattggctg aggttcctgg 240
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gtctagaagt ctcaggacca atgcagcaaa gtctaggagc cctggccaga gctttctggg 600
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<210> 9
<211> 83
<212> PRT
<213> Rattus novartis

<400> 9

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20 25 30
Asn Thr Thr Ala Ile Leu Gln Gly Lys Tyr Gly Leu Leu Leu His Arg
35 40 45
Gly Pro Val His Glu Thr Trp Glu Asp Pro Arg Thr Cys Trp Ile Leu
50 55 60
Cys Val Tyr Tyr Arg Leu Ala Lys Pro His Trp Leu Arg Phe Leu Glu
65 70 75 80
Leu His Leu

<210> 10
<211> 28
<212> DNA
<213> Mus musculus

<400> 10

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<210> 11
<211> 26
<212> DNA
<213> Mus musculus

<400> 11

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